LC-CDR3 having the amino acid sequence of SEQ ID NO:43; or

(h)

a VH region having the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:53

HC-CDR2 having the amino acid sequence of SEQ ID NO:34

HC-CDR3 having the amino acid sequence of SEQ ID NO:35; and

a VL region having the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:58

LC-CDR3 having the amino acid sequence of SEQ ID NO:43; or

(i)

a VH region having the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:33

HC-CDR2 having the amino acid sequence of SEQ ID NO:34

HC-CDR3 having the amino acid sequence of SEQ ID NO:35; and

a VL region having the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:67

LC-CDR3 having the amino acid sequence of SEQ ID NO:43; or

(j)

a VH region having the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:33

HC-CDR2 having the amino acid sequence of SEQ ID NO:34

HC-CDR3 having the amino acid sequence of SEQ ID NO:35; and

a VL region having the following CDRs:

LC-CDR1 having the amino acid sequence of SEQ ID NO:41

LC-CDR2 having the amino acid sequence of SEQ ID NO:42

LC-CDR3 having the amino acid sequence of SEQ ID NO:43.

38. The antigen-binding molecule according to claim **35**, wherein the antigen-binding molecule comprises:

(a) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:310; or

(b) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:297; or

(c) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:294; or

(d) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:299; or

(e) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:301; or

(f) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:289; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:302; or

(g) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:285; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:287; or

(h) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:52; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:57; or

(i) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:62; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:66; or

(j) a VH region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:32; and a VL region comprising an amino acid sequence having at least 70% sequence identity to the amino acid sequence of SEQ ID NO:40.

39. A nucleic acid, or a plurality of nucleic acids, encoding an antigen-binding molecule comprising a heavy chain variable (VH) region and a light chain variable (VL) region, wherein the antigen-binding molecule specifically binds to human VISTA and mouse VISTA, and wherein the antigen-binding molecule inhibits VISTA-mediated signalling independently of Fc-mediated function.

40. The nucleic acid or plurality of nucleic acids according to claim **39**, wherein the antigen-binding molecule comprises:

a VH region incorporating the following CDRs:

HC-CDR1 having the amino acid sequence of SEQ ID NO:305

HC-CDR2 having the amino acid sequence of SEQ ID NO:306

HC-CDR3 having the amino acid sequence of SEQ ID NO:307; and